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TECH CENTER 1600/2300

1600

## RAW SEQUENCE LISTING

DATE: 09/26/2003

PATENT APPLICATION: US/09/772,116A

TIME: 16:02:22

Input Set : A:\Seqlist.txt

Output Set: N:\CRF4\09262003\I772116A.raw

3 <110> APPLICANT: Benjamin, Howard  
4 Findeis, Mark A.  
5 Gefter, Malcolm L.  
6 Musso, Gary  
7 Signer, Ethan R.  
9 <120> TITLE OF INVENTION: METHODS FOR IDENTIFYING COMPOUNDS THAT BIND TO A TARGET  
11 <130> FILE REFERENCE: PPI-010CN  
13 <140> CURRENT APPLICATION NUMBER: US 09/772116A  
14 <141> CURRENT FILING DATE: 2001-01-26  
16 <150> PRIOR APPLICATION NUMBER: US 08/573786  
17 <151> PRIOR FILING DATE: 1995-12-18  
19 <160> NUMBER OF SEQ ID NOS: 6  
21 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
23 <210> SEQ ID NO: 1  
24 <211> LENGTH: 16  
25 <212> TYPE: PRT  
26 <213> ORGANISM: Artificial Sequence  
28 <220> FEATURE:  
29 <223> OTHER INFORMATION: motif  
31 <220> FEATURE:  
32 <221> NAME/KEY: VARIANT  
33 <222> LOCATION: 1, 6, 11, 16  
34 <223> OTHER INFORMATION: Xaa = any amino acid  
36 <220> FEATURE:  
37 <221> NAME/KEY: VARIANT  
38 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15  
39 <223> OTHER INFORMATION: Xaa = Ala or Gly  
41 <400> SEQUENCE: 1

W--> 42 Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa  
43 1 5 10 15

46 <210> SEQ ID NO: 2  
47 <211> LENGTH: 16  
48 <212> TYPE: PRT  
49 <213> ORGANISM: Artificial Sequence  
51 <220> FEATURE:  
52 <223> OTHER INFORMATION: motif  
54 <220> FEATURE:  
55 <221> NAME/KEY: VARIANT  
56 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15  
57 <223> OTHER INFORMATION: Xaa = Ala or Gly  
59 <400> SEQUENCE: 2

W--> 60 Ser Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Met  
61 1 5 10 15

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64 <210> SEQ ID NO: 3  
 65 <211> LENGTH: 16  
 66 <212> TYPE: PRT  
 67 <213> ORGANISM: Artificial Sequence  
 69 <220> FEATURE:  
 70 <223> OTHER INFORMATION: motif  
 72 <220> FEATURE:  
 73 <221> NAME/KEY: VARIANT  
 74 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15  
 75 <223> OTHER INFORMATION: Xaa = Ala or Gly  
 77 <400> SEQUENCE: 3

W--&gt; 78 Ser Xaa Xaa Xaa Xaa Lys Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Gln

79 1 5 10 15

82 <210> SEQ ID NO: 4  
 83 <211> LENGTH: 16  
 84 <212> TYPE: PRT  
 85 <213> ORGANISM: Artificial Sequence  
 87 <220> FEATURE:  
 88 <223> OTHER INFORMATION: motif  
 90 <220> FEATURE:  
 91 <221> NAME/KEY: VARIANT  
 92 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15  
 93 <223> OTHER INFORMATION: Xaa = Ala or Gly  
 95 <400> SEQUENCE: 4

W--&gt; 96 Phe Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Thr

97 1 5 10 15

100 <210> SEQ ID NO: 5  
 101 <211> LENGTH: 16  
 102 <212> TYPE: PRT  
 103 <213> ORGANISM: Artificial Sequence  
 105 <220> FEATURE:  
 106 <223> OTHER INFORMATION: motif  
 108 <220> FEATURE:  
 109 <221> NAME/KEY: VARIANT  
 110 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15  
 111 <223> OTHER INFORMATION: Xaa = Ala or Gly  
 113 <400> SEQUENCE: 5

W--&gt; 114 Ser Xaa Xaa Xaa Xaa Asn Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Ile

115 1 5 10 15

118 <210> SEQ ID NO: 6  
 119 <211> LENGTH: 16  
 120 <212> TYPE: PRT  
 121 <213> ORGANISM: Artificial Sequence  
 123 <220> FEATURE:  
 124 <223> OTHER INFORMATION: motif  
 126 <220> FEATURE:  
 127 <221> NAME/KEY: VARIANT  
 128 <222> LOCATION: 2, 3, 4, 5, 7, 8, 9, 10, 12, 13, 14, 15  
 129 <223> OTHER INFORMATION: Xaa = Ala or Gly

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131 &lt;220&gt; FEATURE:

132 &lt;221&gt; NAME/KEY: VARIANT

133 &lt;222&gt; LOCATION: 16

134 &lt;223&gt; OTHER INFORMATION: Xaa = any amino acid

136 &lt;400&gt; SEQUENCE: 6

W--&gt; 137 Ser Xaa Xaa Xaa Xaa Arg Xaa Xaa Xaa Xaa Leu Xaa Xaa Xaa Xaa Xaa

138 1

5

10

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/772,116A

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 1,2,3,4,5,6,7,8,9,10,11,12,13,14,15,16  
Seq#:2; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15  
Seq#:3; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15  
Seq#:4; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15  
Seq#:5; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15  
Seq#:6; Xaa Pos. 2,3,4,5,7,8,9,10,12,13,14,15,16